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RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/557,796

DATE: 10/20/2000  
TIME: 11:48:15

Input Set : A:\557796.txt  
Output Set: N:\CRF3\10202000\I557796.raw

4 <110> APPLICANT: Hoch, James  
5 Dartois, Veronique  
7 <120> TITLE OF INVENTION: METABOLIC SELECTION METHODS  
10 <130> FILE REFERENCE: 234/191  
12 <140> CURRENT APPLICATION NUMBER: 09/557,796  
13 <141> CURRENT FILING DATE: 2000-04-25  
15 <150> PRIOR APPLICATION NUMBER: 09/172,952  
16 <151> PRIOR FILING DATE: 1998-10-14  
18 <160> NUMBER OF SEQ ID NOS: 33  
20 <170> SOFTWARE: FastSEQ for Windows Version 3.0  
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23 <211> LENGTH: 816  
24 <212> TYPE: DNA  
25 <213> ORGANISM: yia j  
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31 ggcgcattgtt cggtttttttt gggactttttt gggactttttt aaaatgtttttt ttcatcggtt attacagggg 180  
32 ctgcgtttttt ggggtttttt ggcgtttttt ggcgtttttt ggcgtttttt ggcgtttttt ggcgtttttt ggcgtttttt 240  
33 aaatgtttttt ggggtttttt gggactttttt gggactttttt gggactttttt gggactttttt gggactttttt gggactttttt 300  
34 cggcatctttt ggggtttttt gggactttttt gggactttttt gggactttttt gggactttttt gggactttttt gggactttttt 360  
35 gatgaccacg cgtttttttt gggactttttt gggactttttt gggactttttt gggactttttt gggactttttt gggactttttt 420  
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39 gagegaaata tggcgatggaa cggggaaatggaa aacgagctggaa ggcgtttttt gggactttttt gggactttttt gggactttttt 660  
40 cccgtttttt gggactttttt gggactttttt gggactttttt gggactttttt gggactttttt gggactttttt gggactttttt 720  
41 cgccctcaaggc aggtggggaa gaaaatggaaatggaa ctcggatggaa tacggatggaa ggcagaggcg 780  
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63 atgggtttttt gggactttttt gggactttttt gggactttttt gggactttttt gggactttttt gggactttttt gggactttttt 720  
64 tccaaacggat cgtttttttt gggactttttt gggactttttt gggactttttt gggactttttt gggactttttt gggactttttt 780

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128	gccccactct	caggctggat	tatcacgggtt	ctcgactggc	gttggctgtt	tattatcgaa	540
129	ggtttgtct	cgctgggtt	tctgggtctg	tgggcataca	ccatctatga	ccgtccgcag	600
130	gaaggcgcgt	ggatttccga	agcagagaag	cgctatctgg	tcgagacgt	ggccgcggag	660
131	caaaaagcca	ttggccggac	cgaggtaaaa	aacgcctctc	tgagcgcgt	tctctccgac	720
132	aaaaccatgt	ggcagcttat	cgccctgaac	ttcttcttacc	agacccggat	ttacggctac	780
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134	atgtttggca	ttctgcccata	cgtccggcgc	attgtggaa	tgttccctgt	ttctccctt	900
135	tcagaccgaa	ccgttaaaacg	caagctgttc	gtctgctgtc	cgctgattgg	tttcgccttg	960
136	tgcatgttcc	tgtcggtgcc	gtcgaaaaaaac	caaatttggc	tctccatgc	cgcgctggc	1020
137	ggctcgggat	tcttcctgca	atcgccggct	ggcattgttct	ggaccatccc	ggcacgtctg	1080
138	ttcagcgcgg	aaatggccgg	cgccgcgcgc	ggggttatca	acgcgttgg	caacccgcgc	1140
139	ggattttgtg	gcccattatgc	gttcgggggtg	ctgatcacgt	tgtacageaa	agacgcgtggc	1200
140	gtctattggc	tggcgatctc	cctggcgctg	gcccgcgtga	tggcgctgt	gtcgccggcg	1260
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153	tcgccccctgc	cgggttacgc	cgaacgcgcac	atgcgcacgc	tctggcaaca	ctgcgcggcg	180
154	accatttgcg	ggcttattaca	cgaggcgagg	gtatccggcg	aacagattaa	aggcggtggc	240
155	atctccgtctc	agggtcaagg	gcttttctc	ctcgatcaac	aggatcgccg	gtcggttac	300
156	gccatctct	cctccgatcg	tcggggcgtg	aaaatcggtt	agcgctggca	gcccggaccgt	360
157	atccccgaac	ggctctatcc	cggttaccgc	cagacgtgt	ggaccggaca	tccggcttct	420
158	ttgtcgccgt	gggtaaaaaa	gaatgaaccc	cagcgctacg	cgccaaatgg	ctgcgtgatg	480
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162	tgcggggaga	tcaccgtca	ggcagccgt	ttaaccggtc	tggcgccgg	tactcccg	720
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173	gctcggtcg	gtaccggcgt	ataccgcac	tttagcgaag	ccggcgcgc	ccggcagcac	1380
174	ccggcgcga	cgctgtgc	cgatatgacc	gcccacgcgc	gtatcg	caaataccgc	1440
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188 accgaaggc ttacgcgcgt taaaggccctg cgcgcggcgt gtccggggaa gatcatgc  
189 gccactgga aatgcgcgca cgcgcgtgaa accctggcgc agcaggccctt tggcgctgc  
190 gccaactgga tgaccatcat ttgcgcgcga cgcgcgtgaa cggtagagaa aggccacgc  
191 gtggcccgagg cttgcgcggg taaaatttcg atggagctgt tcgcaactg gacgttgat  
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193 gcccaggcca gccccggcaga gtggggggag gcgatctgg cgcgcgtgaa agcgttgcc  
194 gatattggcc tttagatc gattacggc ggcattaccc cagccgtatc acccgctgtt  
195 aaagatataca acgtcaaaacg cttttttgcg gggcgcgcgc tggcaggcgc cggccatccg  
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202 <213> ORGANISM: yia r  
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208 accatgtaaac gccttcgcgc cctggagtgg accccggccc aegcgcgcatac gctgggtgac  
209 gcatgttggaa aaccgcgcgtt cgcattcccgat tcgatgtgtc tgccgcgttca tcgcgttcc  
210 cccttggca gcccgcgtatca aeggtacgc gatcggggcgc gagagattat gaccaaaagcc  
211 atccgcgttggccgcgcgtatca gggatccgc accatccacgc tggcgggttgc cagctgttca  
212 taacgcgttggccgcgcgtatca caccggcgcgcgcgcgcgcgcgcgcgcgcgcgcgcgcgcgc  
213 gaaacggccgc  
214 aactccatca gcaaatggaa aaagtgggac gagatgtttt cgtcaccgtg gtttaccgtc  
215 taccggacgc tccgcgcactt cagcgcgttgg gaaacgcgcgcgcgcgcgcgcgcgcgcgc  
216 ggcacatgcgtatcgc  
217 cttggcccgatc tccgc  
218 aagacgttgc gcgagctgaa ctaccgcgtt tcatttttga ttgagatgtg gacggagaaaa  
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236 tacttctatg ggcgcgtatcc cttgttccgcgcgcgcgcgcgcgcgcgcgcgcgcgcgc  
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238 ggcacatgcgttcc cggcggttattt ggtccatca caccggccctt tggcggggggaa aagacgc  
239 gccacgc  
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240 cggcagtggc cacagctgcc ggatatgcag tctgaactgc tcgataaaca ctatctgcgt 660  
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 255 20 25 30  
 257 Asn Tyr Pro Asn Gly Cys Pro Val Ala His Leu Ser Glu Leu Ala Gly  
 258 35 40 45  
 260 Leu Asn Lys Ser Thr Val His Arg Leu Leu Gln Gly Leu Gln Ser Cys  
 261 50 55 60  
 263 Gly Tyr Val Thr Pro Ala Ala Gly Ser Tyr Ala Leu Thr Thr  
 264 65 70 75 80  
 266 Lys Phe Ile Arg Val Gly Gln Lys Ala Leu Ser Ser Leu Asn Ile Ile  
 267 85 90 95  
 269 His Val Ala Ala Pro His Leu Glu Ala Leu Asn Leu Ala Thr Gly Glu  
 270 100 105 110  
 272 Thr Val Asn Phe Ser Ser Arg Glu Asp Asp His Ala Ile Leu Ile Tyr  
 273 115 120 125  
 275 Lys Leu Glu Pro Thr Thr Gly Met Leu Arg Thr Arg Ala Tyr Ile Gly  
 276 130 135 140  
 278 Gln His Met Arg Cys Thr Ala Arg Gln Trp Ala Lys Ile Tyr Met Ala  
 279 145 150 155 160  
 281 Phe Gly His Pro Asp Tyr Val Glu Ser Tyr Trp Asn Ser His Gln Glu  
 282 165 170 175  
 284 Ile Ile Gln Pro Leu Thr Arg Asn Thr Ile Thr Gly Leu Pro Ala Met  
 285 180 185 190  
 287 His Asp Glu Leu Ala Gln Ile Arg Glu Arg Asn Met Ala Met Asp Arg  
 288 195 200 205  
 289 Glu Glu Asn Glu Leu Gly Val Ser Cys Leu Ala Val Pro Val Phe Asp  
 290 210 215 220  
 292 Ile His Gly Arg Val Pro Tyr Ala Ile Ser Ile Ser Leu Ser Thr Ser  
 293 225 230 235 240  
 295 Arg Leu Lys Gln Val Gly Glu Lys Asn Leu Leu Lys Pro Leu Arg Asp  
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VERIFICATION SUMMARY DATE: 10/20/2000  
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